16 Page 1 of 7

1646

OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/896,096A DATE: 11/23/2001 TIME: 11:07:25

Input Set: N:\Crf3\RULE60\09896096A.TXT Output Set: N:\CRF3\11232001\1896096A.raw

```
<110> APPLICANT: ASHKENAZI, AVI J
         BOTSTEIN, DAVID
 4
 5
         DODGE, KELLY H.
         GURNEY, AUSTIN L.
 6
                                                                ENTERED
 7
         KIM, KYUNG JIN
 8
         LAWRENCE, DAVID A.
 9
         PITTI, ROBERT
         ROY, MARGARET A
10
         TUMAS, DANIEL B
11
12
         WOOD, WILLIAM I.
14 <120> TITLE OF INVENTION: DCR3 Polypeptide, A TNFR Homolog
16 <130> FILE REFERENCE: P1134R2 REVISED
18 <140> CURRENT APPLICATION NUMBER: 09/896,096A
19 <141> CURRENT FILING DATE: 2001-06-28
21 <150> PRIOR APPLICATION NUMBER: US 09/157,289
22 <151> PRIOR FILING DATE: 1998-09-18
24 <150> PRIOR APPLICATION NUMBER: US 60/059,288
25 <151> PRIOR FILING DATE: 1997-09-18
27 <150> PRIOR APPLICATION NUMBER: US 60/094,640
28 <151> PRIOR FILING DATE: 1998-07-30
30 <160> NUMBER OF SEQ ID NOS: 18
32 <210> SEQ ID NO: 1
33 <211> LENGTH: 300
34 <212> TYPE: PRT
35 <213> ORGANISM: Homo sapiens
37 <400> SEQUENCE: 1
38
   Met Arg Ala Leu Glu Gly Pro Gly Leu Ser Leu Leu Cys Leu Val
39
41
    Leu Ala Leu Pro Ala Leu Leu Pro Val Pro Ala Val Arg Gly Val
42
44
   Ala Glu Thr Pro Thr Tyr Pro Trp Arg Asp Ala Glu Thr Gly Glu
45
47
   Arg Leu Val Cys Ala Gln Cys Pro Pro Gly Thr Phe Val Gln Arg
48
                     50
                                                               60
50
    Pro Cys Arg Arg Asp Ser Pro Thr Thr Cys Gly Pro Cys Pro Pro
51
                                                               75
                     65
53
   Arg His Tyr Thr Gln Phe Trp Asn Tyr Leu Glu Arg Cys Arg Tyr
54
                                                               90
                     80
                                          85
   Cys Asn Val Leu Cys Gly Glu Arg Glu Glu Glu Ala Arg Ala Cys
56
57
                                         100
                     95
59
   His Ala Thr His Asn Arg Ala Cys Arg Cys Arg Thr Gly Phe Phe
60
                    110
                                         115
                                                              120
62
   Ala His Ala Gly Phe Cys Leu Glu His Ala Ser Cys Pro Pro Gly
                                         130
65
   Ala Gly Val Ile Ala Pro Gly Thr Pro Ser Gln Asn Thr Gln Cys
66
                    140
                                         145
                                                              150
   Gln Pro Cys Pro Pro Gly Thr Phe Ser Ala Ser Ser Ser Ser Ser
68
```

Input Set: N:\Crf3\RULE60\09896096A.TXT
Output Set: N:\CRF3\11232001\1896096A.raw

```
69
                    155
                                         160
                                                              165
71
    Glu Gln Cys Gln Pro His Arg Asn Cys Thr Ala Leu Gly Leu Ala
72
                    170
74
    Leu Asn Val Pro Gly Ser Ser Ser His Asp Thr Leu Cys Thr Ser
75
                    185
                                         190
                                                              195
77
    Cys Thr Gly Phe Pro Leu Ser Thr Arg Val Pro Gly Ala Glu Glu
78
                                         205
                                                              210
                    200
80
    Cys Glu Arq Ala Val Ile Asp Phe Val Ala Phe Gln Asp Ile Ser
81
                    215
                                         220
                                                              225
    Ile Lys Arg Leu Gln Arg Leu Gln Ala Leu Glu Ala Pro Glu
83
84
                    230
                                         235
                                                              240
86
    Gly Trp Gly Pro Thr Pro Arg Ala Gly Arg Ala Ala Leu Gln Leu
                                                              255
87
                    245
                                         250
89
    Lys Leu Arg Arg Arg Leu Thr Glu Leu Leu Gly Ala Gln Asp Gly
90
                    260
                                                              270
92
    Ala Leu Leu Val Arg Leu Leu Gln Ala Leu Arg Val Ala Arg Met
93
                    275
                                         280
                                                              285
95
    Pro Gly Leu Glu Arg Ser Val Arg Glu Arg Phe Leu Pro Val His
96
                    290
                                         295
98 <210> SEQ ID NO: 2
99 <211> LENGTH: 1114
100 <212> TYPE: DNA
101 <213> ORGANISM: Homo sapiens
103 <220> FEATURE:
104 <221> NAME/KEY: Unsure
105 <222> LOCATION: 1090
106 <223> OTHER INFORMATION: Unknown base
108 <400> SEQUENCE: 2
    tccgcaggcg gaccgggggc aaaggaggtg gcatgtcggt caggcacagc 50
111
     agggteetgt gteegegetg ageegegete teeetgetee ageaaggaee 100
113
     atgagggcgc tggagggcc aggcctgtcg ctgctgtgcc tggtgttggc 150
115
     getgeetgee etgetgeegg tgeeggetgt acgeggagtg geagaaacae 200
117
     ccacctaccc ctggcgggac gcagagacag gggagcggct ggtgtgcgcc 250
119
     cagtgeecec caggeacett tgtgcagegg cegtgeegee gagacageee 300
121 cacqacqtqt qqcccqtqtc caccqcqcca ctacacqcaq ttctqqaact 350
123
     acctggagcg ctgccgctac tgcaacqtcc tctgcgggga gcgtgaggag 400
     gaggeacggg cttgccacgc cacccacaac cgtgcctgcc gctgccgcac 450
127
     cggcttcttc gcgcacgctg gtttctgctt ggagcacgca tcgtgtccac 500
129
     ctggtgccgg cgtgattgcc ccgggcaccc ccagccagaa cacgcagtgc 550
131
     caqceqtqcc ccccaqqcac cttctcaqcc aqcaqctcca qctcaqaqca 600
133
     gtgccagccc caccgcaact gcacggccct gggcctggcc ctcaatgtgc 650
135
     caggetette etcecatgae accetgtgea ceagetgeae tggetteece 700
137
     ctcagcacca gggtaccagg agctgaggag tgtgagcgtg ccgtcatcga 750
139
     ctttgtggct ttccaggaca tctccatcaa gaggctgcag cggctgctgc 800
141
     aggeeetega ggeeeggag ggetggggte egacaceaag ggegggeege 850
143
     geggeettge agetgaaget gegteggegg etcaeggage teetggggge 900
145
     gcaggacggg gcgctgctgg tgcggctgct gcaggcgctg cgcgtggcca 950
     ggatgcccgg gctggagcgg agcgtccgtg agcgcttcct ccctgtgcac 1000
147
149
     tgatcctggc cccctcttat ttattctaca tccttggcac cccacttgca 1050
```

Input Set : N:\Crf3\RULE60\09896096A.TXT
Output Set: N:\CRF3\11232001\1896096A.raw

W--> 151 ctgaaagagg cttttttta aatagaagaa atgaggtttn ttaaaaaaaa 1100 153 aa'aaaaaaaa aaaa 1114 155 <210> SEQ ID NO: 3 156 <211> LENGTH: 491 157 <212> TYPE: DNA 158 <213> ORGANISM: Unknown 160 <220> FEATURE: 161 <223> OTHER INFORMATION: Unknown organism 163 <220> FEATURE: 164 <221> NAME/KEY: unsure 165 <222> LOCATION: 62, 73, 86, 98 166 <223> OTHER INFORMATION: unknown base 168 <400> SEQUENCE: 3 169 gccqaqacaq ccccacqacq tqtqqcccqt qtccaccqcq ccactacacq 50 W--> 171 cagttctgga antaactgga genetgeege tactgnaacg teetctgngg 100 ggagcgtgag gaggaggcac gggcttgcca cgccacccac aaccgtgcct 150 173 175 qccqctqccq caccqqcttc ttcqcqcacq ctqqtttctq cttqqaqcac 200 177 qcatcgtgtc cacctggtgc cggcgtgatt gccccgggca cccccaqcca 250 179 gaacacgcag tgcctagccg tgccccccag gcaccttctc agccagcagc 300 181 tccagctcag agcagtgcca gccccaccgc aactgcacgg ccctgggcct 350 183 ggccctcaat gtgccaggct cttcctccca tgacaccctg tgcaccagct 400 185 gcactggctt cccctcagc accagggtac caggagctga ggagtgtgag 450 187 cgtgccgtca tcgactttgt ggctttccag gacatctcca t 491 189 <210> SEQ ID NO: 4 190 <211> LENGTH: 73 191 <212> TYPE: DNA 192 <213> ORGANISM: Unknown 194 <220> FEATURE: 195 <223> OTHER INFORMATION: Unknown organism 197 <400> SEQUENCE: 4 198 gccqaqacag ccccacgacg tgtggcccgt gtccaccgcg ccactacacg 50 200 cattetggaa etacetggag ege 73 202 <210> SEO ID NO: 5 203 <211> LENGTH: 271 204 <212> TYPE: DNA 205 <213> ORGANISM: Unknown 207 <220> FEATURE: 208 <223> OTHER INFORMATION: Unknown organism 210 <220> FEATURE: 211 <221> NAME/KEY: unsure 212 <222> LOCATION: 42, 62, 73, 86, 98, 106, 120, 122, 153, 167, 184, 220, 233 213 <223> OTHER INFORMATION: unknown base 215 <400> SEQUENCE: 5 W--> 216 gccgagacag ccccacgacg tgtggcccgt gtccaccgcg cnactacacg 50 W--> 218 cagttctgga antaactgga gcnctgccgc tactgnaacg tcctctgngg 100 W--> 220 ggagentgag gaggaggean gngettgeea egecaceeae aacegegeet 150 W--> 222 gengetgeag caceggntte ttegegeaeg etgntttetg ettggageae 200 W--> 224 gcatcgtgtc cacctggtgn cggcgtgatt gcnccgggca cccccagcca 250 gaacacgcat gcaaagccgt g 271

Input Set : N:\Crf3\RULE60\09896096A.TXT
Output Set: N:\CRF3\11232001\1896096A.raw

228 <210> SEQ ID NO: 6 229 <211> LENGTH: 201 230 <212> TYPE: DNA 231 <213> ORGANISM: Unknown 233 <220> FEATURE: 234 <223> OTHER INFORMATION: Unknown organism 236 <220> FEATURE: 237 <221> NAME/KEY: unsure 238 <222> LOCATION: 182 239 <223> OTHER INFORMATION: unknown base 241 <400> SEQUENCE: 6 242 gcagttctgg aactacctgg agcgctgccg ctactgcaac gtcctctgcg 50 244 gggagcgtga ggaggaggca cgggcttgcc acgccaccca caaccgtgcc 100 246 tgccgctgcc gcaccggctt cttcgcgcac gctggtttct gcttggagca 150 W--> 248 cgcatcgtgt ccacctggtg ccggcgtgat tnccccgggc acccccagcc 200 250 a 201 252 <210> SEQ ID NO: 7 253 <211> LENGTH: 277 254 <212> TYPE: DNA 255 <213> ORGANISM: Unknown 257 <220> FEATURE: 258 <223> OTHER INFORMATION: Unknown organism 260 <220> FEATURE: 261 <221> NAME/KEY: unsure 262 <222> LOCATION: 142 · 263 <223> OTHER INFORMATION: unknown base 265 <400> SEQUENCE: 7 266 gaggggcccc caggagtggt ggccggaggt gtggcagggg tcaggttgct 50 268 ggtcccagcc ttgcaccctg agctaggaca ccagttcccc tgaccctgtt 100 W--> 270 cttccctcct ggctgcaggc acccccagcc agaacacgca gnccagccgt 150 272 gccccccagg caccttctca gccagcagct ccagctcaga gcagtgccag 200 274 ccccaccgca actgcacggc cctgggcctg gccctcaatg tgccaggctc 250 276 ttcctcccat gacaccctgt gcaccag 277 278 <210> SEQ ID NO: 8 279 <211> LENGTH: 199 280 <212> TYPE: DNA 281 <213> ORGANISM: Unknown 283 <220> FEATURE: 284 <223> OTHER INFORMATION: Unknown organism 286 <400> SEQUENCE: 8 gcatcgtgtc cacctggtgc cggcgtgatt gccccgggca cccccagcca 50 289 gaacacgcag gcctagccgt gcccccagg caccttctca gccagcagct 100 291 ccagctcaga gcagtgccag ccccaccgca actgcacggc cctgggcctg 150 293 gccctcaatg tgccaggctc ttcctcccat gacaccctgt gcaccagct 199 295 <210> SEQ ID NO: 9 296 <211> LENGTH: 226 297 <212> TYPE: DNA 298 <213> ORGANISM: Unknown

300 <220> FEATURE:

Input Set: N:\Crf3\RULE60\09896096A.TXT
Output Set: N:\CRF3\11232001\1896096A.raw

301 <223> OTHER INFORMATION: Unknown organism 303 <220> FEATURE: 304 <221> NAME/KEY: unsure 305 <222> LOCATION: 4, 9, 12, 165 306 <223> OTHER INFORMATION: unknown base 308 <400> SEQUENCE: 9 W--> 309 agengtgene encaggeace tteteageea geagtteeag eteagageag 50 311 tgccagcccc accgcaactg cacggccctg ggcctggccc tcaatgtgcc 100 313 aggetettee teecatgaca egetgtgeac eagetgeact ggetteeece 150 W--> 315 tcagcaccag ggtancagga gctgaggagt gtgagcgtgc cgtcatcgac 200 317 tttgtggctt tccaggacat ctccat 226 319 <210> SEQ ID NO: 10 320 <211> LENGTH: 283 321 <212> TYPE: DNA 322 <213> ORGANISM: Homo sapiens 324 <220> FEATURE: 325 <221> NAME/KEY: Unsure 326 <222> LOCATION: 1-283 327 <223> OTHER INFORMATION: Unknown organism 329 <220> FEATURE: 330 <221> NAME/KEY: unsure 331 <222> LOCATION: 27, 64, 140 332 <223> OTHER INFORMATION: unknown base 334 <400> SEQUENCE: 10 W--> 335 cttgtccacc tggtgccggc gtgattnccc gggcaccccc agccagaaca 50 W--> 337 egeagtgeea geenteecce caggeacett etcagecage agetecaget 100 W--> 339 cagagcagtg ccagccccac cgcaactgca acgccctggn ctggccctca 150 341 atgtgccagg ctcttcctcc catgacaccc tgtgcaccag ctgcactggc 200 343 ttccccctca gcaccagggt accaggagct gaggagtgtg agcgtgccgt 250 345 catcgacttt gtggctttcc aggacatctc cat 283 347 <210> SEQ ID NO: 11 348 <211> LENGTH: 21 349 <212> TYPE: DNA 350 <213> ORGANISM: Unknown 352 <220> FEATURE: 353 <223> OTHER INFORMATION: Unknown organism 355 <400> SEQUENCE: 11 356 cacgctggtt tctgcttgga g 21 358 <210> SEQ ID NO: 12 359 <211> LENGTH: 22 360 <212> TYPE: DNA 361 <213> ORGANISM: Unknown 363 <220> FEATURE: 364 <223> OTHER INFORMATION: Unknown organism 366 <400> SEQUENCE: 12 367 agctggtgca cagggtgtca tg 22 369 <210> SEQ ID NO: 13 370 <211> LENGTH: 53

371 <212> TYPE: DNA

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/896,096A

DATE: 11/23/2001 TIME: 11:07:26

Input Set : N:\Crf3\RULE60\09896096A.TXT
Output Set: N:\CRF3\11232001\1896096A.raw

```
L:151 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:171 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:216 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:218 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:220 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:222 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:224 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:248 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:248 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:270 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:309 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:315 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:335 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:337 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:339 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
```